

# NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES

Applicant must file the items indicated below within the time period set the Office action to which the Notice is attached to avoid abandonment under 35 U.S.C. § 133 (extensions of time may be obtained under the provisions of 37 CFR 1.136(a)).

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 C.F.R. 1.821 - 1.825 for the following reason(s):

- ☐ 1. This application clearly fails to comply with the requirements of 37 C.F.R. 1.821-1.825. Applicant's attention is directed to the final rulemaking notice published at 55 FR 18230 (May 1, 1990), and 1114 OG 29 (May 15, 1990). If the effective filing date is on or after July 1, 1998, see the final rulemaking notice published at 63 FR 29620 (June 1, 1998) and 1211 OG 82 (June 23, 1998).
- ☐ 2. This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 C.F.R. 1.821(c).
- ☐ 3. A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 C.F.R. 1.821(e).
- ☒ 4. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 C.F.R. 1.822 and/or 1.823, as indicated on the attached copy of the marked -up "Raw Sequence Listing."
- ☐ 5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A Substitute computer readable form must be submitted as required by 37 C.F.R. 1.825(d).
- ☐ 6. The paper copy of the "Sequence Listing" is not the same as the computer readable form of the "Sequence Listing" as required by 37 C.F.R. 1.821(e).
- ☐ 7. Other: \_

## Applicant Must Provide:

- ☒ An initial or substitute computer readable form (CRF) copy of the "Sequence Listing".
- ☒ An initial or substitute paper copy of the "Sequence Listing", as well as an amendment directing its entry into the specification.
- ☒ A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 C.F.R. 1.821(e) or 1.821(f) or 1.821(g).

For questions regarding compliance to these requirements, please contact:

- For Rules Interpretation, call (703) 308-4216 or (703) 308-2923
- For CRF Submission Help, call (703) 308-4212
- For PatentIn software Program Support:
  - HELP DESK: (703) 739-8559, ext 508, M-F, 8 AM to 5 PM EST except holidays
  - Email: [PATIN21HELP@uspto.gov](mailto:PATIN21HELP@uspto.gov)
  - To purchase PatentIn software: (703) 306-2600

PLEASE RETURN A COPY OF THIS NOTICE WITH YOUR RESPONSE

Gabriele E. Bugaisky

AU 1653

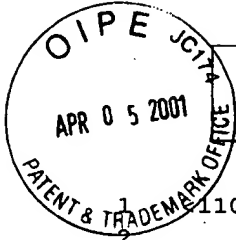
3/15/01 (703)308-4201

p#10

PAGE: 1

RAW SEQUENCE LISTING  
PATENT APPLICATION, US/09/493,795DATE: 02/17/2000  
TIME: 12:51:55

Input Set: I493795.RAW



This Raw Listing contains the General Information  
Section and up to first 5 pages.

PP. 5, 2-3

Does Not Comply  
Corrected Diskette Needed

110> APPLICANT: Watkins, Maren  
Olivera, Baldomero M.  
Hillyard, David R.  
McIntosh, J. Michael  
Jones, Robert M.  
<120> TITLE OF INVENTION: Alpha-Conotoxin Peptides  
<130> FILE REFERENCE: Alphas 2  
<140> CURRENT APPLICATION NUMBER: US/09/493,795  
<141> CURRENT FILING DATE: 2000-01-28  
<150> EARLIER APPLICATION NUMBER: US 60/118,381  
<151> EARLIER FILING DATE: 1999-01-29  
<160> NUMBER OF SEQ ID NOS: 404  
<170> SOFTWARE: PatentIn Ver. 2.0  
<210> SEQ ID NO 1  
<211> LENGTH: 17  
<212> TYPE: PRT  
<213> ORGANISM: Artificial Sequence  
<220> FEATURE:  
<223> OTHER INFORMATION: Description of Artificial Sequence: Alpha-Conotoxin  
Peptide Generic Formula I  
<220> FEATURE:  
<221> NAME/KEY: PEPTIDE  
<222> LOCATION: (1)..(3)  
<223> OTHER INFORMATION: Xaa at residue 1 is des-Xaa, Ile, Leu or Val; Xaa  
at residue 2 is des-Xaa, Ala or Gly; Xaa at  
residue 3 is des-Xaa, Gly, Trp (D or L), neo-Trp,  
halo-Trp or any unnatural aromatic amino acid.  
<220> FEATURE:  
<221> NAME/KEY: PEPTIDE  
<222> LOCATION: (4)..(5)  
<223> OTHER INFORMATION: Xaa at residue 4 is des-Xaa, Gly, Trp (D or L),  
neo-Trp, halo-Trp or any unnatural aromatic amino  
acid; Xaa at residue 5 is Glu, gamma-carboxy-Glu  
(Gla), Asp, Ala, Thr, Ser, Gly, Ile, Tyr, nor-Tyr,  
<220> FEATURE:  
<221> NAME/KEY: PEPTIDE  
<222> LOCATION: (5)..(8)  
<223> OTHER INFORMATION: mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr,  
O-phospho-Tyr, nitro-Tyr or any unnatural hydroxy  
containing amino acid; Xaa at residue 8 is Ser,  
Thr, Arg, ornithine, homoarginine, Lys,  
N-methyl-Lys,  
<220> FEATURE:  
<221> NAME/KEY: PEPTIDE

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45 <222> LOCATION: (8)..(9)  
46 <223> OTHER INFORMATION: N,N-dimethyl-Lys, N,N,N-trimethyl-Lys or any  
47 unnatural basic amino acid; Xaa at residue 9 is  
48 Asp, Glu, Gla, Arg, ornithine, homoarginine, Lys,  
49 N-methyl-Lys, N,N-dimethyl-Lys,  
50 N,N,N-trimethyl-Lys or  
51 <220> FEATURE:  
52 <221> NAME/KEY: PEPTIDE  
53 <222> LOCATION: (9)..(11)  
54 <223> OTHER INFORMATION: any unnatural basic amino acid; Xaa at residue 10  
55 is Ser, Thr, Asn, Ala, Gly, His, halo-His, Pro or  
56 hydroxy-Pro; Xaa at residue 11 is Thr, Ser, Ala,  
57 Asp, Asn, Pro, hydroxy-Pro,  
58 <220> FEATURE:  
59 <221> NAME/KEY: PEPTIDE  
60 <222> LOCATION: (11)..(13)  
61 <223> OTHER INFORMATION: Arg, ornithine, homoarginine, Lys, N-methyl-Lys,  
62 N,N-dimethyl-Lys, N,N,N-trimethyl-Lys or any  
63 unnatural basic amino acid; Xaa at residue 13 is  
64 Gly, Ser, Thr, Ala, Asn,  
65 <220> FEATURE:  
66 <221> NAME/KEY: PEPTIDE  
67 <222> LOCATION: (13)..(14)  
68 <223> OTHER INFORMATION: Arg, ornithine, homoarginine, Lys, N-methyl-Lys,  
69 N,N-dimethyl-Lys, N,N,N-trimethyl-Lys or any  
70 unnatural basic amino acid; Xaa at residue 14 is  
71 Gln, Leu, His, halo-His, Trp (D or L), halo-Trp,  
72 neo-Trp,  
73 <220> FEATURE:  
74 <221> NAME/KEY: PEPTIDE  
75 <222> LOCATION: (14)  
76 <223> OTHER INFORMATION: Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr,  
77 O-sulpho-Tyr, O-phospho-Tyr, nitro-Tyr, Arg,  
78 ornithine, homoarginine, Lys, N-methyl-Lys,  
79 N,N-dimethyl-Lys, N,N,N-trimethyl-Lys, any  
80 unnatural basic amino  
81 <220> FEATURE:  
82 <221> NAME/KEY: PEPTIDE  
83 <222> LOCATION: (14)..(15)  
84 <223> OTHER INFORMATION: acid or any unnatural aromatic amino acid; Xaa at  
85 residue 15 is Asn, His, halo-His, Ile, Leu, Val,  
86 Gln, Arg, ornithine, homoarginine, Lys,  
87 N-methyl-Lys, N,N-dimethyl-Lys,  
88 N,N,N-trimethyl-Lys or any  
89 <220> FEATURE:  
90 <221> NAME/KEY: PEPTIDE  
91 <222> LOCATION: (15)..(16)  
92 <223> OTHER INFORMATION: unnatural basic amino acid; Xaa at residue 16 is  
93 des-Xaa, Val, Ile, Leu, Arg, ornithine,  
94 homoarginine, Lys, N-methyl-Lys, N,N-dimethyl-Lys,

Cys is at residue 16 - do y.  
mean  
17  
Xaa  
is at  
location

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RAW SEQUENCE LISTING  
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95 N,N,N-trimethyl-Lys or any unnatural basic amino  
96 acid.

97 <400> SEQUENCE: 1  
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99 1 5 10 15  
W--> 100 Xaa

101 <210> SEQ ID NO 2  
102 <211> LENGTH: 21  
103 <212> TYPE: PRT  
104 <213> ORGANISM: Artificial Sequence  
105 <220> FEATURE:  
106 <223> OTHER INFORMATION: Description of Artificial Sequence: Alpha-Conotoxin  
107 Peptide Generic Formula II.  
108 <220> FEATURE:  
109 <221> NAME/KEY: PEPTIDE  
110 <222> LOCATION: (1)..(3)  
111 <223> OTHER INFORMATION: Xaa at residue 1 is des-Xaa, Asp, Glu or  
112 gamma-carboxy-Glu (Gla); Xaa at residue 2 is  
113 des-Xaa, Gln, Ala, Asp, Glu, Gla; Xaa at residue 3  
114 is des-Xaa, Gly, Ala, Asp, Glu, Gla, Pro or  
115 hydroxy-Pro.  
116 <220> FEATURE:  
117 <221> NAME/KEY: PEPTIDE  
118 <222> LOCATION: (4)..(7)  
119 <223> OTHER INFORMATION: Xaa at residue 4 is des-Xaa4, Gly, Glu, Gla, Gln,  
120 Asp, Asn, Pro or hydroxy-Pro; Xaa at residue 7 is  
121 Ser, Thr, Gly, Glu, Gla, Asn, Trp (D or L),  
122 neo-Trp, halo-Trp, Arg, ornithine, homoarginine,  
123 <220> FEATURE:  
124 <221> NAME/KEY: PEPTIDE  
125 <222> LOCATION: (7)  
126 <223> OTHER INFORMATION: Lys, N-methyl-Lys, N,N-dimethyl-Lys,  
127 N,N,N-trimethyl-Lys, any unnatural basic amino  
128 acid, Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr,  
129 O-sulpho-Tyr, O-phospho-Tyr, nitro-Tyr or any  
130 unnatural hydroxy  
131 <220> FEATURE:  
132 <221> NAME/KEY: PEPTIDE  
133 <222> LOCATION: (7)..(8)  
134 <223> OTHER INFORMATION: containing amino acid; Xaa at residue 8 is Asp,  
135 Asn, His, halo-His, Thr, Ser, Tyr, nor-Tyr,  
136 mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr,  
137 O-phospho-Tyr, nitro-Tyr or any unnatural hydroxy  
138 <220> FEATURE:  
139 <221> NAME/KEY: PEPTIDE  
140 <222> LOCATION: (8)..(10)  
141 <223> OTHER INFORMATION: containing amino acid; Xaa at residue 9 is Pro or  
142 hydroxy-Pro; Xaa at residue 10 is Ala, Ser, Thr,  
143 Asp, Val, Ile, Pro, hydroxy-Pro, Tyr, nor-Tyr,  
144 mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr,

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145 <220> FEATURE:  
146 <221> NAME/KEY: PEPTIDE  
147 <222> LOCATION: (10)..(12)  
148 <223> OTHER INFORMATION: O-phospho-Tyr, nitro-Tyr or any unnatural hydroxy  
149 containing amino acid; Xaa at residue 12 is Gly,  
150 Ile, Leu, Val, Ala, Thr, Ser, Pro, hydroxy-Pro,  
151 Phe, Trp (D or L), neo-Trp, halo-Trp, Arg,  
152 ornithine,  
153 <220> FEATURE:  
154 <221> NAME/KEY: PEPTIDE  
155 <222> LOCATION: (12)..(13)  
156 <223> OTHER INFORMATION: homoarginine, Lys, N-methyl-Lys, N,N-dimethyl-Lys,  
157 N,N,N-trimethyl-Lys, any unnatural basic amino  
158 acid or any unnatural aromatic amino acid; Xaa at  
159 residue 13 is Ala, Asn, Phe, Pro, hydroxy-Pro,  
160 <220> FEATURE:  
161 <221> NAME/KEY: PEPTIDE  
162 <222> LOCATION: (13)  
163 <223> OTHER INFORMATION: Glu, Gla, Gln, His, halo-His, Val, Ser, Thr, Arg,  
164 ornithine, homoarginine, Lys, N-methyl-Lys,  
165 N,N-dimethyl-Lys, N,N,N-trimethyl-Lys or any  
166 unnatural basic amino acid.  
167 <220> FEATURE:  
168 <221> NAME/KEY: PEPTIDE  
169 <222> LOCATION: (14)  
170 <223> OTHER INFORMATION: Xaa at residue 14 is Thr, Ser, His, halo-His, Leu,  
171 Ile, Val, Asn, Met, Pro, hydroxy-Pro, Arg,  
172 ornithine, homoarginine, Lys, N-methyl-Lys,  
173 N,N-dimethyl-Lys, N,N,N-trimethyl-Lys, any  
174 unnatural basic  
175 <220> FEATURE:  
176 <221> NAME/KEY: PEPTIDE  
177 <222> LOCATION: (14)..(15)  
178 <223> OTHER INFORMATION: amino acid, Tyr, nor-Tyr, mono-halo-Tyr,  
179 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr,  
180 nitro-Tyr or any unnatural hydroxy containing  
181 amino acid; Xaa at residue 15 is Asn, Pro,  
182 hydroxy-Pro, Gln, Ser, Thr,  
183 <220> FEATURE:  
184 <221> NAME/KEY: PEPTIDE  
185 <222> LOCATION: (15)  
186 <223> OTHER INFORMATION: Arg, ornithine, homoarginine, Lys, N-methyl-Lys,  
187 N,N-dimethyl-Lys N,N,N-trimethyl-Lys, any  
188 unnatural basic amino acid, Tyr, nor-Tyr,  
189 mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr,  
190 O-phospho-Tyr, nitro-Tyr  
191 <220> FEATURE:  
192 <221> NAME/KEY: PEPTIDE  
193 <222> LOCATION: (15)..(16)  
194 <223> OTHER INFORMATION: or any unnatural hydroxy containing amino acid;

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# RAW SEQUENCE LISTING PATENT APPLICATION US/09/493,795

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195 Xaa at residue 16 is des-Xaa, Gly, Thr, Ser, Pro,  
196 hydroxy-Pro, Tyr, nor-Tyr, mono-halo-Tyr,  
197 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr,  
198 nitro-Tyr or any  
199 <220> FEATURE:  
200 <221> NAME/KEY: PEPTIDE  
201 <222> LOCATION: (16)..(17)  
202 <223> OTHER INFORMATION: unnatural hydroxy containing amino acid; Xaa at  
203 residue 17 is des-Xaa14, Ile, Val, Asp, Leu, Phe,  
204 Arg, ornithine, homoarginine, Lys, N-methyl-Lys,  
205 N,N-dimethyl-Lys, N,N,N-trimethyl-Lys, any  
206 unnatural  
207 <220> FEATURE:  
208 <221> NAME/KEY: PEPTIDE  
209 <222> LOCATION: (17)..(19)  
210 <223> OTHER INFORMATION: basic amino acid, Tyr, nor-Tyr, mono-halo-Tyr,  
211 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr,  
212 nitro-Tyr or any unnatural hydroxy containing  
213 amino acid; Xaa at residue 19 is des-Xaa, Gly,  
214 Ala, Met, Ser,  
215 <220> FEATURE:  
216 <221> NAME/KEY: PEPTIDE  
217 <222> LOCATION: (19)  
218 <223> OTHER INFORMATION: Thr, Trp (D or L), neo-Trp, halo-Trp, any  
219 unnatural aromatic amino acid, Arg, ornithine,  
220 homoarginine, Lys, N-methyl-Lys, N,N-dimethyl-Lys,  
221 N,N,N-trimethyl-Lys or any unnatural basic amino  
222 acid.  
223 <220> FEATURE:  
224 <221> NAME/KEY: PEPTIDE  
225 <222> LOCATION: (20)  
226 <223> OTHER INFORMATION: Xaa at residue 20 is des-Xaa, Trp (D or L),  
227 neo-Trp, halo-Trp, any unnatural aromatic amino  
228 acid, Arg, ornithine, homoarginine, Lys,  
229 N-methyl-Lys, N,N-dimethyl-Lys,  
230 N,N,N-trimethyl-Lys or any  
231 <220> FEATURE:  
232 <221> NAME/KEY: PEPTIDE  
233 <222> LOCATION: (20)..(21)  
234 <223> OTHER INFORMATION: unnatural basic amino acid; Xaa at residue 21 is  
235 des-Xaa, Arg, ornithine, homoarginine, Lys,  
236 N-methyl-Lys, N,N-dimethyl-Lys,  
237 N,N,N-trimethyl-Lys or any unnatural basic amino  
238 acid.  
239 <400> SEQUENCE: 2  
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241 1 5 10 15  
W--> 242 Xaa Cys Xaa Xaa Xaa  
243 20

&lt;210&gt; SEQ ID NO 3

**Please Note:**

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

PSI

Input Set: I493795.RAW

Line	Error/Warning	Original Text
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100	W "N" or "Xaa" used: Feature required	Xaa
240	W "N" or "Xaa" used: Feature required	Xaa Xaa Xaa Xaa Cys Cys Xaa Xaa Xaa Xaa C
242	W "N" or "Xaa" used: Feature required	Xaa Cys Xaa Xaa Xaa
459	W "N" or "Xaa" used: Feature required	Xaa Xaa Xaa Xaa Xaa Cys Cys Xaa Xaa Xaa X
461	W "N" or "Xaa" used: Feature required	Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa X
474	W "N" or "Xaa" used: Feature required	Asp Xaa Cys Cys Ser Asp Ser Arg Cys Gly X
485	W "N" or "Xaa" used: Feature required	Ala Cys Cys Ser Asp Arg Arg Cys Arg Xaa A
505	W "N" or "Xaa" used: Feature required	Asp Xaa Cys Cys Arg Arg His Ala Cys Thr L
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577	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser Asp Xaa Arg Cys Xaa His G
590	W "N" or "Xaa" used: Feature required	Cys Cys Ser Asp Xaa Ala Cys Xaa Gln Thr X
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655	W "N" or "Xaa" used: Feature required	Xaa Xaa Cys Cys Asn Asn Xaa Ala Cys Val X
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682	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser Arg Xaa Xaa Cys Ala Val L
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733	W "N" or "Xaa" used: Feature required	Xaa Gln Cys Cys Ser His Xaa Ala Cys Asn V
747	W "N" or "Xaa" used: Feature required	Xaa Xaa Cys Cys Ser His Xaa Ala Cys Asn V
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785	W "N" or "Xaa" used: Feature required	Xaa Gln Cys Cys Ser His Xaa Ala Cys Asn V
798	W "N" or "Xaa" used: Feature required	Xaa Gln Cys Cys Ser His Xaa Ala Cys Asn V
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824	W "N" or "Xaa" used: Feature required	Xaa Gln Cys Cys Ser His Xaa Ala Cys Asn I
837	W "N" or "Xaa" used: Feature required	Xaa Gln Cys Cys Ser His Xaa Ala Cys Asn V
851	W "N" or "Xaa" used: Feature required	Gly Gly Cys Cys Ser His Xaa Ala Cys Ala V
864	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Ala Cys Ser Val A
876	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Ala Cys Asn Val A
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892	W "N" or "Xaa" used: Feature required	Arg Xaa Ser
903	W "N" or "Xaa" used: Feature required	Xaa Cys Cys Ser His Xaa Ala Cys Ser Gly A
916	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Ala Cys Ser Gly A
931	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Xaa Cys Ala Met A
945	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Xaa Cys Phe Leu A
958	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser Asn Xaa Xaa Cys Ile Ala X
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981	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser Asn Xaa Ala Cys Ala Gly A
993	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser Arg Xaa Ala Cys Ile Ala A
1006	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser Asn Xaa Val Cys His Val X
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1037	W "N" or "Xaa" used: Feature required	Xaa Xaa Cys Cys Ser Asp Xaa Arg Cys Asn S

Input Set: I493795.RAW

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1050	W "N" or "Xaa" used: Feature required	Xaa Gln Cys Cys Ser Asp Xaa Arg Cys Asn V
1064	W "N" or "Xaa" used: Feature required	Xaa Val Cys Cys Ser Asp Xaa Arg Cys Asn V
1076	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser Arg Xaa Xaa Cys Ile Ala A
1088	W "N" or "Xaa" used: Feature required	Xaa Gln Cys Cys Ser His Leu Ala Cys Asn V
1108	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser Xaa Phe Asp Cys Arg Met M
1110	W "N" or "Xaa" used: Feature required	Gly Xaa Arg
1129	W "N" or "Xaa" used: Feature required	Gly Gly Cys Cys Ser Phe Ala Ala Cys Arg X
1145	W "N" or "Xaa" used: Feature required	Gly Gly Cys Cys Phe His Xaa Val Cys Xaa I
1166	W "N" or "Xaa" used: Feature required	Ser Ala Thr Cys Cys Asn Xaa Xaa Xaa Cys X
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2845	W "N" or "Xaa" used: Feature required	Gly Cys Cys Xaa Asn Xaa Val Cys Xaa Xaa X
2863	W "N" or "Xaa" used: Feature required	Xaa Xaa Xaa Gly Cys Cys Arg His Xaa Ala C
2874	W "N" or "Xaa" used: Feature required	Cys Cys Ala Asp Xaa Asp Cys Arg Phe Arg X
2895	W "N" or "Xaa" used: Feature required	Gly Cys Cys Xaa Asn Xaa Ser Cys Xaa Xaa X
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2916	W "N" or "Xaa" used: Feature required	Cys Cys Ser Asn Xaa Thr Cys Xaa Xaa Thr X
2929	W "N" or "Xaa" used: Feature required	Cys Cys Ala Asn Xaa Ile Cys Xaa Asn Thr X
2948	W "N" or "Xaa" used: Feature required	Cys Cys Asn Asn Xaa Thr Cys Xaa Xaa Thr X
2968	W "N" or "Xaa" used: Feature required	Cys Cys Ser Asn Xaa Val Cys Xaa Xaa Thr X
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3018	W "N" or "Xaa" used: Feature required	Gly Cys Cys Xaa Asn Xaa Xaa Cys Gly Ala S
3032	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser Xaa Xaa Xaa Cys Phe Ala T
3046	W "N" or "Xaa" used: Feature required	Gly Gly Cys Cys Ser Xaa Xaa Xaa Cys Ile A
3061	W "N" or "Xaa" used: Feature required	Gly Gly Cys Cys Ser Xaa Xaa Xaa Cys Ile A
3073	W "N" or "Xaa" used: Feature required	Asp Cys Cys Ser Asn Xaa Xaa Cys Ser Gln A
3084	W "N" or "Xaa" used: Feature required	Asp Cys Cys Ser Asn Xaa Xaa Cys Ala His A
3097	W "N" or "Xaa" used: Feature required	Xaa Cys Cys Thr Asn Xaa Val Cys His Ala X
3111	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser Asn Xaa Val Cys His Leu X
3124	W "N" or "Xaa" used: Feature required	Xaa Cys Cys Thr Asn Xaa Val Cys His Val X
3145	W "N" or "Xaa" used: Feature required	Xaa Xaa Cys Cys Ser Xaa Xaa Ala Cys Asn L
3159	W "N" or "Xaa" used: Feature required	Xaa Xaa Cys Cys Ser Asp Xaa Arg Cys Asn S
3181	W "N" or "Xaa" used: Feature required	Leu Asn Cys Cys Met Ile Xaa Xaa Cys Xaa X
3183	W "N" or "Xaa" used: Feature required	Cys Ser Xaa Val Arg
3198	W "N" or "Xaa" used: Feature required	Ala Phe Gly Cys Cys Asp Leu Ile Xaa Cys L
3200	W "N" or "Xaa" used: Feature required	Arg Cys Asn Xaa Val His
3221	W "N" or "Xaa" used: Feature required	Leu Gly Cys Cys Asn Val Thr Xaa Cys Xaa X
3223	W "N" or "Xaa" used: Feature required	Cys Asn Xaa Val Arg
3235	W "N" or "Xaa" used: Feature required	Asp Xaa Cys Cys Ser Asn Xaa Ala Cys Arg V
3258	W "N" or "Xaa" used: Feature required	Leu Asn Cys Cys Ser Ile Xaa Gly Cys Xaa A
3260	W "N" or "Xaa" used: Feature required	Cys Ser Xaa Val Arg
3274	W "N" or "Xaa" used: Feature required	Gly Gly Cys Cys Ser His Xaa Val Cys Xaa P
3286	W "N" or "Xaa" used: Feature required	Gly Gly Cys Cys Ser His Xaa Val Cys Asn L
3301	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Xaa Cys Xaa Ala A
3314	W "N" or "Xaa" used: Feature required	Gly Gly Cys Cys Ser His Xaa Ala Cys Ser V
3330	W "N" or "Xaa" used: Feature required	Gly Gly Cys Cys Ser Xaa Xaa Ala Cys Ser V
3351	W "N" or "Xaa" used: Feature required	Val Ser Cys Cys Val Val Arg Xaa Cys Xaa I
3353	W "N" or "Xaa" used: Feature required	Cys Leu Xaa Ala Asp Xaa Arg Thr Leu
3374	W "N" or "Xaa" used: Feature required	Xaa Asn Cys Cys Ser Ile Xaa Gly Cys Xaa X



Input Set: I493795.RAW

Line	Error/Warning	Original Text
3376	W "N" or "Xaa" used: Feature required	Cys Ser Xaa Val Arg
3388	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser Asn Xaa Val Cys His Leu X
3402	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser Asn Xaa Ile Cys Xaa Phe A
3422	W "N" or "Xaa" used: Feature required	Xaa Cys Cys Ser Gln Xaa Xaa Cys Arg Xaa X
3434	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Ala Cys Ala Gly A
3445	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ala Val Xaa Ser Cys Arg Leu A
3457	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Ala Cys Asn Val A
3470	W "N" or "Xaa" used: Feature required	Thr Xaa Xaa Xaa Cys Cys Xaa Asn Xaa Xaa C
3485	W "N" or "Xaa" used: Feature required	Asp Ala Cys Cys Ser Asp Xaa Arg Cys Ser G
3498	W "N" or "Xaa" used: Feature required	Xaa Asp Cys Cys Ser Asp Xaa Arg Cys Ser V
3510	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Ala Cys Ala Gly S
3522	W "N" or "Xaa" used: Feature required	Xaa Asp Cys Cys Ser Asp Xaa Arg Cys Ser V
3534	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Ala Cys Ala Gly A
3548	W "N" or "Xaa" used: Feature required	Gly Cys Cys Gly Asn Xaa Ser Cys Ser Ile H
3561	W "N" or "Xaa" used: Feature required	Thr Asp Ser Xaa Xaa Cys Cys Leu Asp Ser A
3577	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser Asn Xaa Xaa Cys Xaa Ala A
3589	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Ala Cys Ser Val A
3608	W "N" or "Xaa" used: Feature required	Gly Xaa Cys Cys Ile Asn Asp Ala Cys Arg S
3623	W "N" or "Xaa" used: Feature required	Gly Cys Cys Xaa Asn Ile Ala Cys Arg Ile A
3643	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Val Cys Arg Phe A
3659	W "N" or "Xaa" used: Feature required	Asp Xaa Cys Cys Ala Ser Xaa Xaa Cys Arg L
3679	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser Asn Xaa Val Cys Xaa Gln A
3681	W "N" or "Xaa" used: Feature required	Arg Xaa Ser
3694	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Xaa Cys Ala Gln A
3709	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Ala Cys Ser Gly A
3711	W "N" or "Xaa" used: Feature required	Arg Xaa Ser
3725	W "N" or "Xaa" used: Feature required	Asp Xaa Cys Cys Ser Xaa Xaa Asp Cys Gly A
3745	W "N" or "Xaa" used: Feature required	Xaa Cys Cys Ser Gln Xaa Xaa Cys Arg Xaa X
3757	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Ala Cys Ala Gly A
3771	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser Asp Xaa Ser Cys Asn Val A
3784	W "N" or "Xaa" used: Feature required	Xaa Xaa Cys Cys Ser Asp Xaa Arg Cys Ser V
3797	W "N" or "Xaa" used: Feature required	Gly Gly Cys Cys Ser Asn Xaa Ala Cys Leu V
3809	W "N" or "Xaa" used: Feature required	Arg Asp Xaa Cys Cys Phe Asn Xaa Ala Cys A
3822	W "N" or "Xaa" used: Feature required	Cys Cys Ser Asp Xaa Ser Cys Xaa Arg Leu H
3835	W "N" or "Xaa" used: Feature required	Cys Cys Thr Asn Xaa Ala Cys Leu Val Asn A
3847	W "N" or "Xaa" used: Feature required	Asp Xaa Cys Cys Ser Asp Xaa Arg Cys His G
3859	W "N" or "Xaa" used: Feature required	Asp Cys Cys Ser His Xaa Leu Cys Arg Leu P
3879	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Val Cys Xaa Val A
3891	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Ala Cys Asn Val A
3910	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Val Cys Xaa Val A
3923	W "N" or "Xaa" used: Feature required	Gly Gly Cys Cys Ser His Xaa Ala Cys Xaa V
3936	W "N" or "Xaa" used: Feature required	Val Cys Cys Ser Asn Xaa Val Cys His Val A
3949	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Val Cys Asn Leu S
3963	W "N" or "Xaa" used: Feature required	Xaa Xaa Cys Cys Ser His Xaa Ala Cys Asn V
3975	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser Asn Xaa Ala Cys Leu Val A
3987	W "N" or "Xaa" used: Feature required	Asp Cys Cys Asp Asp Xaa Ala Cys Thr Val A
4001	W "N" or "Xaa" used: Feature required	Gly Cys Cys Ser Asn Xaa Xaa Cys Ile Ala X
4024	W "N" or "Xaa" used: Feature required	Cys Cys Thr Ile Xaa Ser Cys Xaa Xaa Xaa X
4038	W "N" or "Xaa" used: Feature required	Gly Cys Cys Gly Asn Xaa Ala Cys Ser Gly S
4357	W Invalid/Missing Amino Acid Numbering	

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VERIFICATION SUMMARY  
PATENT APPLICATION US/09/493,795

DATE: 02/17/2006  
TIME: 12:51:55

Input Set: I493795.RAW

Line ? Error/Warning

Original Text

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4440 W Invalid/Missing Amino Acid Numbering